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			10		20		30		40
			TPVSEKQLAE	VVANTITPLM	KAQSVPGMAV	AVIYQGKPHY			
50	60	70		80	90	100			
YTFGKADIAA	NKPVTPQTLF	ELGSISKTFT	GVLGGDAIAR	GEISLDDAVT	RYWPQLTGKQ				
		* *							
110	120	130	140	150	160				
WQGIRMLDLA	TYTAGGLPLQ	VPDEVTDNAS	LLRFYQNWQP	QWKPGTTRLY	ANASIGLFGA				
				*	*				
170	180	190	200	210	220				
LAVKPSGMPY	EQAMTTRVLK	PLKLDHTWIN	VPKAEAAHYA	WGYRDGKAVR	VSPGMLDAQA				
230	240	250	260	270	280				
YGVKTNVQDM	ANWVMANMAP	ENVADASLKQ	GIALAQSRYW	RIGSMYQGLG	WEMLNWPVEA				
290	300	310	320	330	340				
NTVVEGSDSK	VALAPLPVAE	VNPPAPPVKA	SWVHKTGSTG	GFGSYVAFIP	EKQIGIVMLA				
			** *						
350	360								
NTSYPNPARV	EAAYHILEAL	Q							

FIGURE 1

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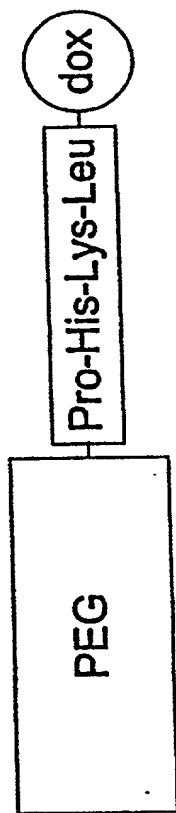


FIGURE 2

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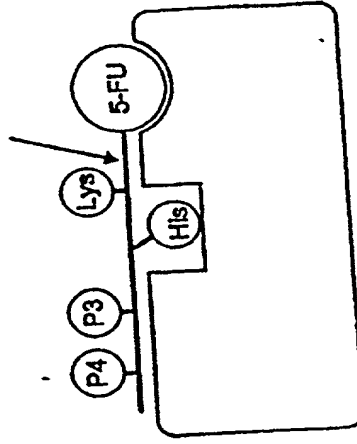


FIGURE 3

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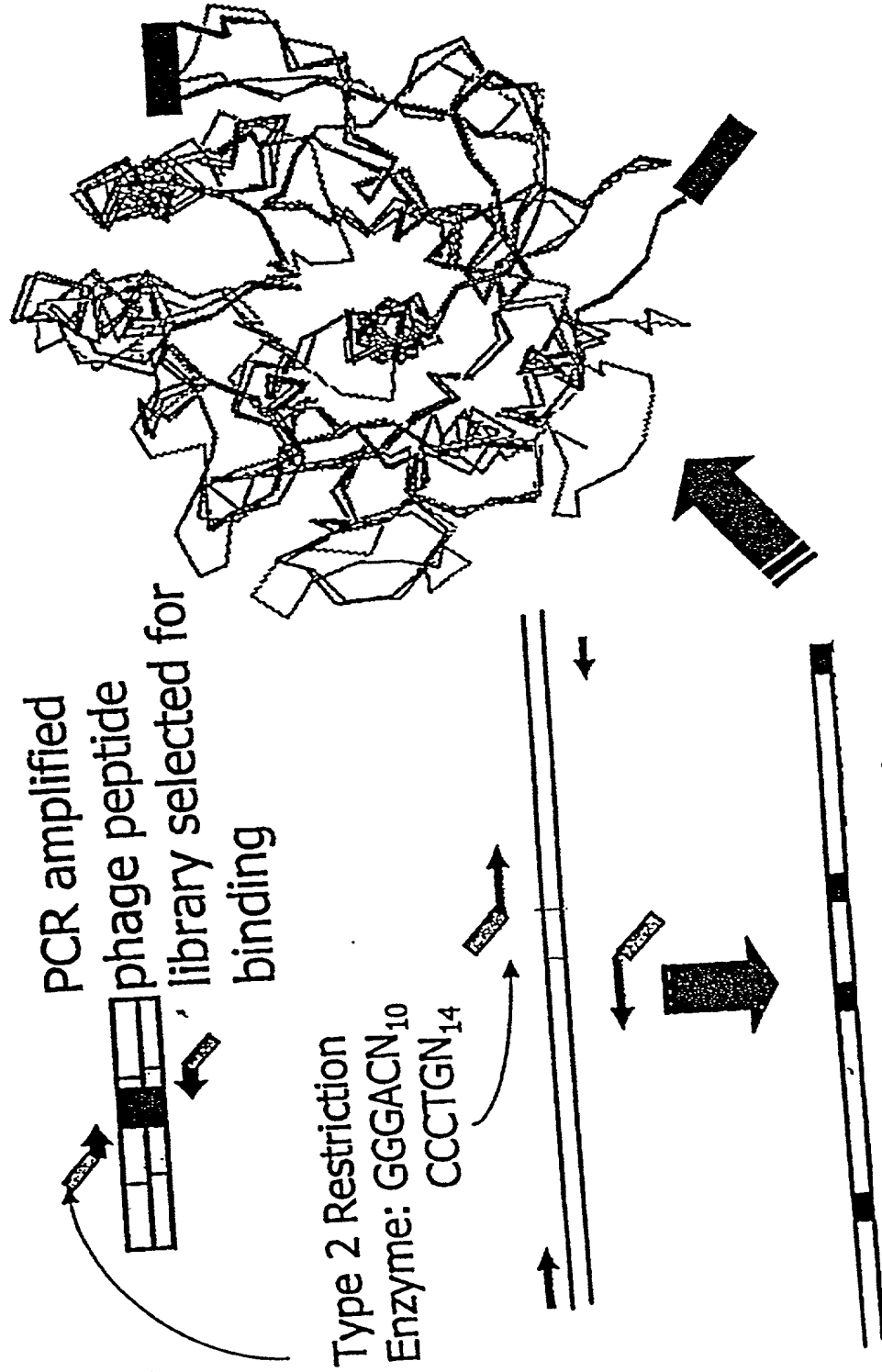


FIGURE 4

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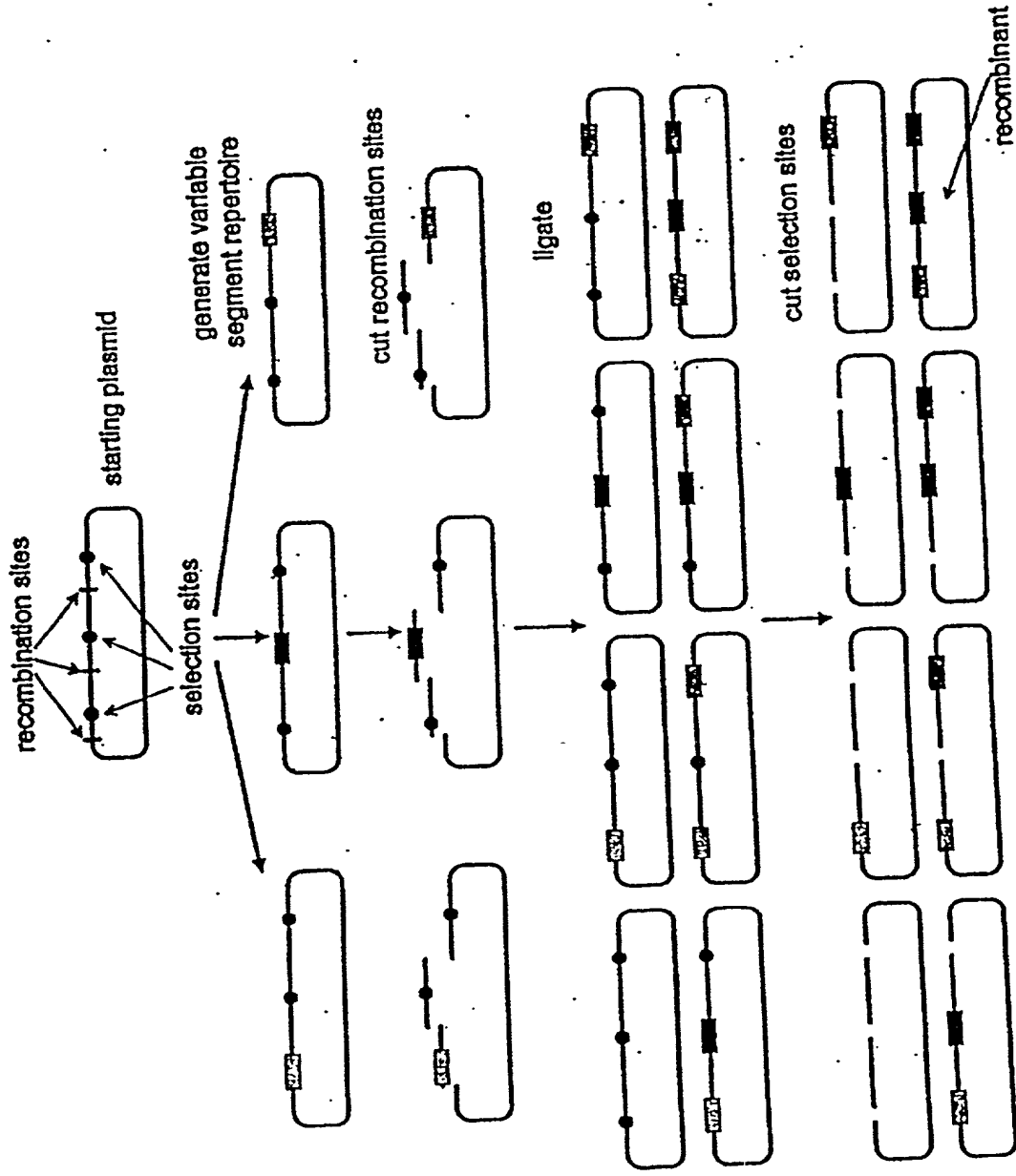


FIGURE 5

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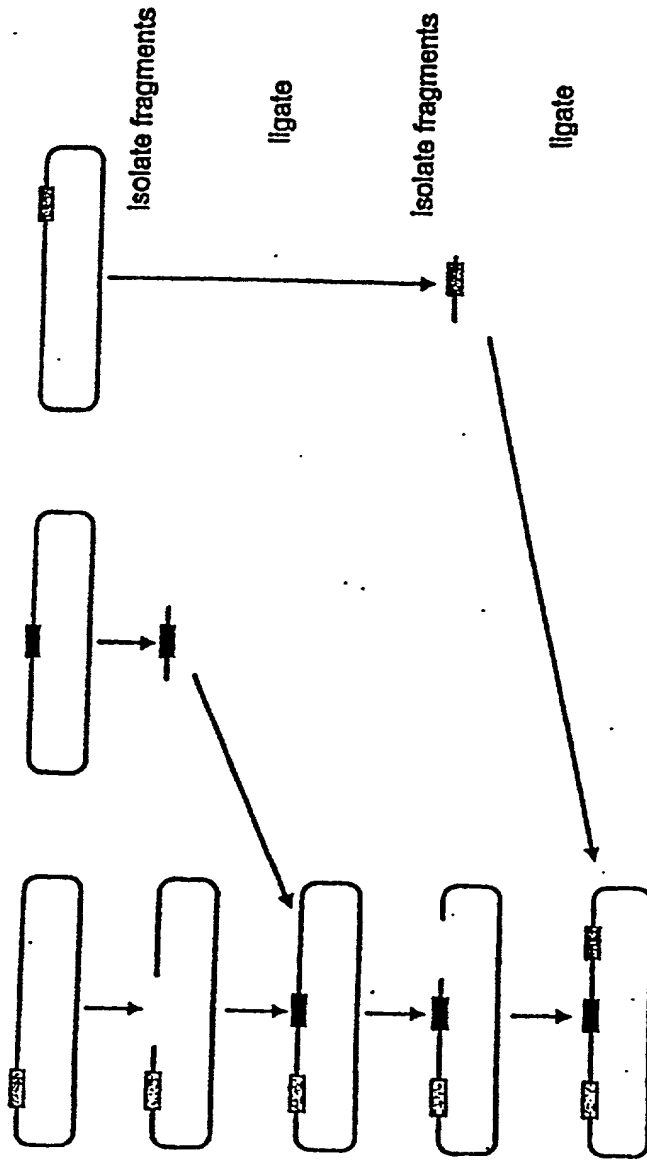


FIGURE 6

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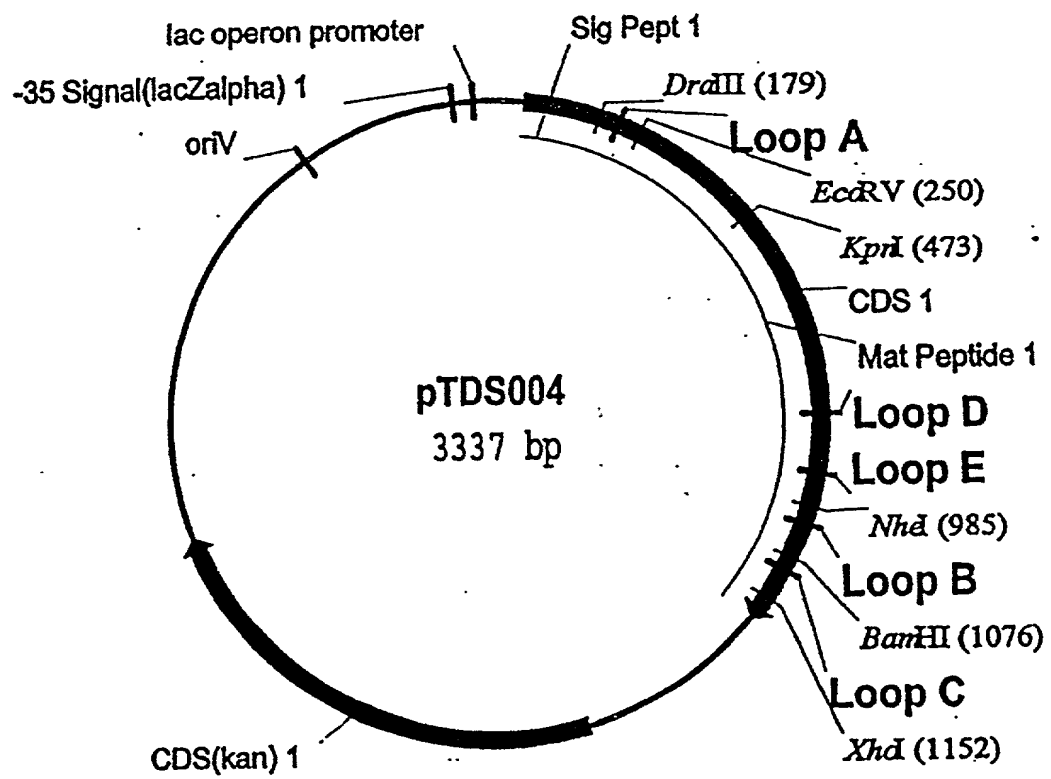


FIGURE 7

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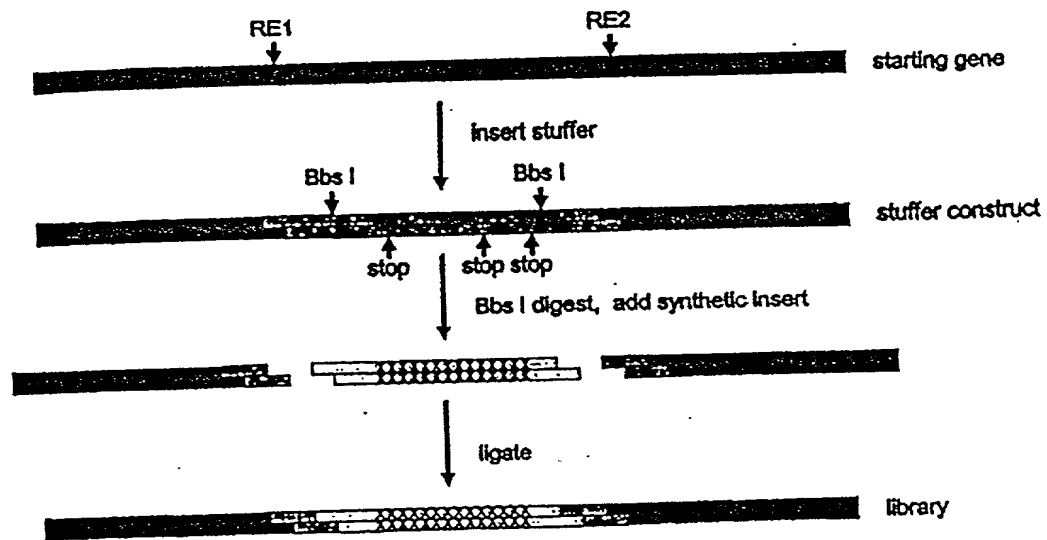


FIGURE 8

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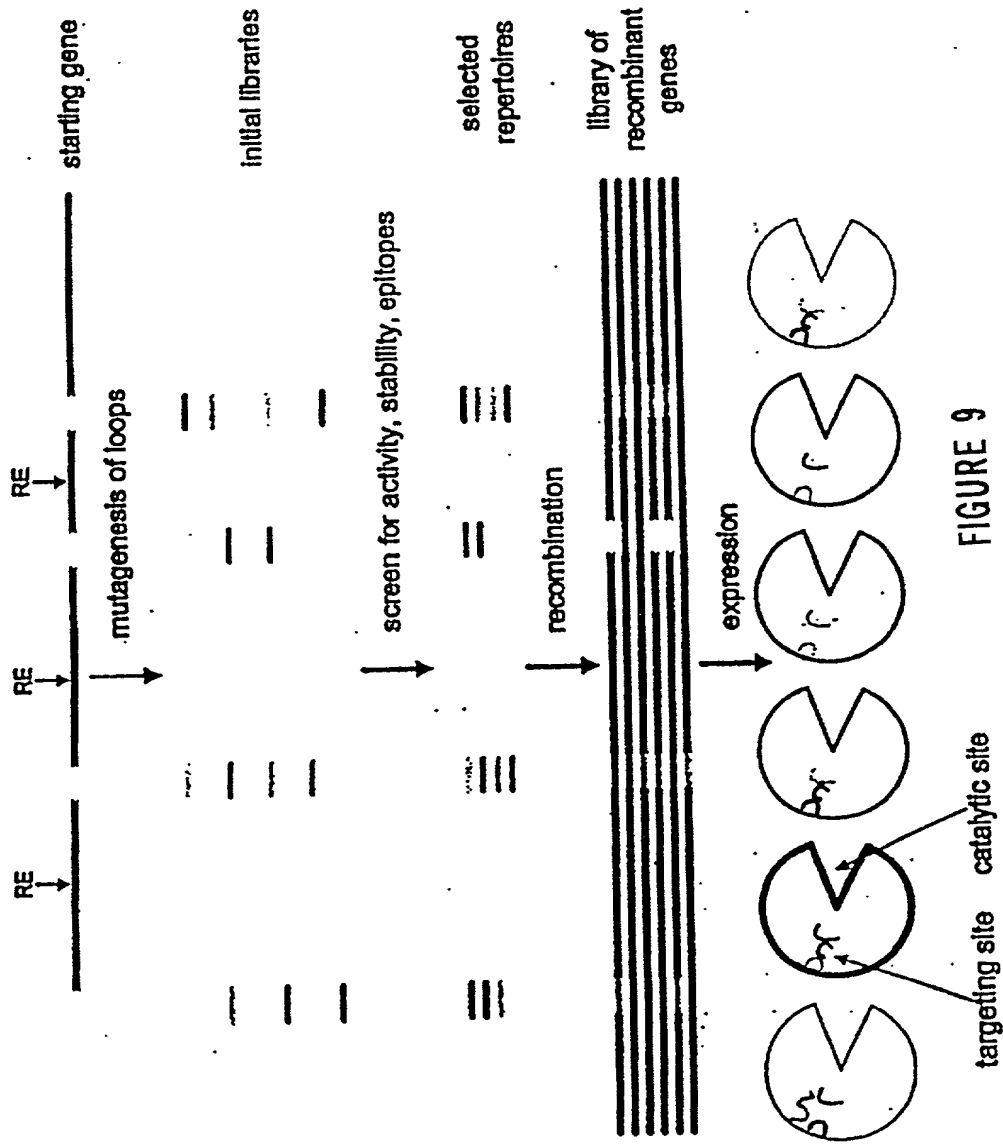


FIGURE 9

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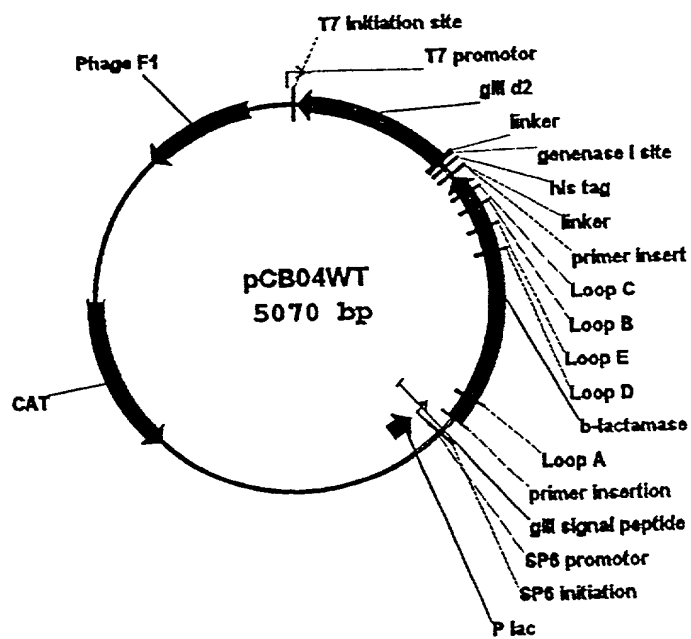


FIGURE 10